

SCORE Search Results Details for Application 10804772 and Search Result us-10-804-772- 1.rng.

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This page gives you Search Results detail for the Application 10804772 and Search Result us-10-804-772-1.rng.

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GenCore version 5.1.9
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OM nucleic - nucleic search, using sw model

Run on: July 31, 2006, 11:41:24 ; Search time 2548 Seconds
(without alignments)
18848.056 Million cell updates/sec

Title: US-10-804-772-1
Perfect score: 6888
Sequence: 1 atgtgggtattatattgttg.....cctagttcaggccaaagctt 6888

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 5244920 seqs, 3486124231 residues

Total number of hits satisfying chosen parameters: 10489840

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : N_Geneseq_8:*

- 1: geneseqn1980s:*
- 2: geneseqn1990s:*
- 3: geneseqn2000s:*
- 4: geneseqn2001as:*
- 5: geneseqn2001bs:*
- 6: geneseqn2002as:*
- 7: geneseqn2002bs:*
- 8: geneseqn2003as:*
- 9: geneseqn2003bs:*
- 10: geneseqn2003cs:*
- 11: geneseqn2003ds:*
- 12: geneseqn2004as:*
- 13: geneseqn2004bs:*
- 14: geneseqn2005s:*
- 15: geneseqn2006s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result	No.	Score	% Query Match	Length	DB	ID	Description
	1	6888	100.0	6888	3	AAA59599	Aaa59599 DNA encod
	2	3379	49.1	3379	14	AED95597	Aed95597 Arabidops
	3	669	9.7	1035	14	AED95588	Aed95588 Cytochrom
c	4	140.4	2.0	8056	8	ABZ10246	Abz10246 Haematopo
	5	136.4	2.0	8056	8	ABZ10246	Abz10246 Haematopo
	6	135.6	2.0	6508	10	ADF31314	Adf31314 DNA encod
	7	132.8	1.9	1509	14	AED95599	Aed95599 Rice puta
	8	132.8	1.9	1521	12	ADK14855	Adk14855 Rice OsDW
	9	127.8	1.9	2066	13	ADX64858	Adx64858 Plant ful
c	10	124	1.8	8056	8	ABZ10100	Abz10100 Haematopo
	11	123.2	1.8	8056	8	ABZ10100	Abz10100 Haematopo
c	12	112.2	1.6	2482	14	ADZ71091	Adz71091 Human chr
	13	107	1.6	2131	14	ADZ71009	Adz71009 Human chr
c	14	107	1.6	2131	14	ADZ71009	Adz71009 Human chr
	15	102.6	1.5	2482	14	ADZ71091	Adz71091 Human chr
	16	94.6	1.4	1181	9	ACH03759	Ach03759 Rice ster
	17	94.2	1.4	1428	14	ADZ70899	Adz70899 Human mat
	18	94.2	1.4	1554	14	ADZ70999	Adz70999 Human chr
	19	93.8	1.4	1524	14	ADZ70897	Adz70897 Human mat
	20	93.8	1.4	4660	14	ADZ70902	Adz70902 Human mat
	21	92.8	1.3	4937	2	AAT85307	Aat85307 Arabidops
c	22	91.6	1.3	960	11	ACN85231	Acn85231 Breast ca
	23	91.2	1.3	828	14	ADZ70929	Adz70929 Human chr
	24	91.2	1.3	1367	14	ADZ70967	Adz70967 Human chr
c	25	91.2	1.3	1428	14	ADZ70899	Adz70899 Human mat
	26	89.8	1.3	115218	8	ACA64845	Aca64845 Human HNR
c	27	89.6	1.3	1554	14	ADZ70999	Adz70999 Human chr
c	28	89.6	1.3	110000	13	ABD32968_6	Continuation (7 of
c	29	89.2	1.3	4660	14	ADZ70902	Adz70902 Human mat
c	30	88.8	1.3	1169	14	ADZ71002	Adz71002 Human chr
	31	88.2	1.3	453	9	ACH03747	Ach03747 Rice ster
c	32	88.2	1.3	5286	13	ADS89278	Ads89278 Oligonucl
c	33	88.2	1.3	5286	13	ADS89552	Ads89552 Oligonucl
	34	87.6	1.3	568	10	ABX56964	Abx56964 Arabidops
	35	87.6	1.3	1419	6	ABZ12814	Abz12814 Arabidops
	36	87.6	1.3	1608	2	AAT85306	Aat85306 Arabidops
	37	87.6	1.3	1646	3	AAC48157	Aac48157 Arabidops
	38	87.6	1.3	1649	3	AAC47986	Aac47986 Arabidops
	39	87.6	1.3	1649	15	AEF30128	Aef30128 Lead_Cere
	40	87.6	1.3	1682	15	AEF09798	Aeg09798 Thale cre
c	41	87.4	1.3	158001	12	ADL17884	Adl17884 Human pho
c	42	86.4	1.3	38678	14	AEB32373	Aeb32373 Human gen
c	43	86.4	1.3	38684	14	AEB32391	Aeb32391 Human gen
	44	86.4	1.3	158001	12	ADL17884	Adl17884 Human pho
	45	86.2	1.3	5286	13	ADS89552	Ads89552 Oligonucl

ALIGNMENTS

RESULT 1
AAA59599